SEQUENCE LISTING

<110>   Zeng, Z et al.

<120>   Human Tumor Necrosis Factor receptors TR21 and TR22

<130>   PF530C1

<140>   To be Assigned
<141>   2003-07-17

<150>   09/910,562
<151>   2001-07-23

<150>   60/221,143
<151>   2000-07-27

<150>   60/220,116
<151>   2000-07-24

<160>   10

<170>   PatentIn version 3.1

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acaatgggag cagccgcaca ttgcactcca gaacagagac gaccccgtcg cccagcaacg   240

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cagagtagca ggtgagccgt ggttttggtg acattggggg cagagtggtg caggggtgagg   960

agaaggtact tggagcctcc caggtgctgt ggcagcatag gaatggtatt tgacagggaa   1020

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        35   40   45

Asn Asp Thr Gly Asn Gly His Pro Glu Tyr Ile Ala Tyr Ala Leu Val
        50   55   60

Pro Val Phe Phe Ile Met Gly Leu Phe Gly Val Leu Ile Cys His Leu
65   70   75   80

Leu Lys Lys Lys Gly Tyr Arg Cys Thr Thr Glu Ala Glu Gln Asp Ile
                85   90   95

Glu Glu Glu Lys Val Glu Lys Ile Glu Leu Asn Asp Ser Val Asn Glu
            100   105   110

Asn Ser Asp Thr Val Gly Gln Ile Val His Tyr Ile Met Lys Asn Glu
        115   120   125

Ala Asn Ala Asp Val Leu Lys Ala Met Val Ala Asp Asn Ser Leu Tyr
        130   135   140

Asp Pro Glu Ser Pro Val Thr Pro Ser Thr Pro Gly Ser Pro Pro Val
145   150   155   160

Ser Pro Gly Pro Leu Ser Pro Gly Gly Thr Pro Gly Lys His Val Cys
                165   170   175

Gly His His Leu His Thr Val Gly Gly Val Val Glu Arg Asp Val Cys
            180   185   190

His Arg Cys Arg His Lys Arg Trp His Phe Ile Lys Pro Thr Asn Lys
            195   200   205

2

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Ser Arg Glu Ser Arg Pro Arg Arg Gln Gly Glu Val Thr Val Leu Ser
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Val Gly Arg Phe Arg Val Thr Lys Val Glu His Lys Ser Asn Gln Lys
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acctccagcc tgcaggacgg agccccctcc catcatcaca cagtgcacct gggctctgca   360

gccccttgcc tccattgcag ccgcagcaag aggcctccac ttgtccgtca gggacgctcc   420

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            20              25              30

Lys Gly Tyr Arg Cys Arg Thr Ser Arg Gly Ser Glu Pro Asp Asp Ala
        35              40              45

Gln Leu Gln Pro Pro Glu Asp Asp Asp Met Asn Glu Asp Thr Val Glu
    50              55              60

Arg Ile Val Arg Cys Ile Ile Gln Asn Glu Ala Asn Ala Glu Ala Leu
65              70              75              80
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3

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Lys Glu Met Leu Gly Asp Ser Glu Gly Glu Gly Thr Val Gln Leu Ser
              85                      90                      95

Ser Val Asp Ala Thr Ser Ser Leu Gln Asp Gly Ala Pro Ser His His
              100                     105                     110

His Thr Val His Leu Gly Ser Ala Ala Pro Cys Leu His Cys Ser Arg
              115                     120                     125

Ser Lys Arg Pro Pro Leu Val Arg Gln Gly Arg Ser Lys Glu Gly Lys
         130                     135                     140

Ser Arg Pro Arg Thr Gly Glu Thr Thr Val Phe Ser Val Gly Arg Phe
145                     150                     155                160

Arg Val Thr His Ile Glu Lys Arg Tyr Gly Leu His Glu His Arg Asp
                   165                     170                     175

Gly Ser Pro Thr Asp Arg Ser Trp Gly Ser Gly Gly Gly Gln Asp Pro
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Gly Val


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<212>   PRT
<213>   homo sapiens

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Ser Thr His Ala Ser Gly Pro Pro Ala Pro Glu Arg Leu Cys Leu Pro
1               5                   10                  15

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            20   25   30

Thr Gln Asn Ile Leu His Thr Arg Leu Ser Leu Cys Ser Leu Ser Trp
        35   40   45

Val Ser Leu Ala Ser Ser Phe Xaa Xaa Leu Ala Xaa Arg Arg Lys Ala
    50   55   60

Ile Val Val Gln Gln Lys Gln Ser Lys Ile Ser Lys Lys Lys Lys Val
65   70   75   80

Glu Lys Xaa Xaa Leu Asn Asp Ser Val Asn Glu Asn Ser Asp Thr Val
            85   90   95

Gly Gln Ile Val His Tyr Ile Met Lys Asn Glu Ala Asn Ala Asp Val
            100   105   110

Leu Lys Ala Met Val Ala Asp Asn Ser Leu Tyr Asp Pro Glu Ser Pro
        115   120   125

Val Thr Pro Ser Thr Pro Gly Ser Pro Pro Val Ser Pro Gly Leu Cys
    130   135   140

His Gln Gly Gly Arg Gln Gly Ser Thr Ser Val Ala Ile Ile Cys Ile
145   150   155   160

Arg Trp Ala Val Xaa Ser Arg Gly Met Cys Val Ile Gly Val Gly Thr
            165   170   175

Ser Gly Gly Thr Leu
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<210>  10
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acaagagcag gtggcagcag gggaacgtct tctcatgctc cgtgatgcat gaggctctgc         660

acaaccacta cacgcagaag agcctctccc tgtctccggg taaatgagtg cgacggccgc         720

gactctagag gat                                                             733
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6